Data Mining:

Concepts and Techniques

(3rd ed.)

- Chapter 10 -

Jiawei Han, Micheline Kamber, and Jian Pei

Chapter 10. Cluster Analysis: Basic Concepts and Methods

- Cluster Analysis: Basic Concepts 💆
- **Partitioning Methods**
- **Hierarchical Methods**
- **Density-Based Methods**
- **Grid-Based Methods**
- **Evaluation of Clustering**
- Summary

What is Cluster Analysis?

- Cluster: A collection of data objects
 - Similar (or related) to one another within the same group
 - Dissimilar (or unrelated) to the objects in other groups
- Cluster analysis (or clustering, data segmentation, ...)
 - Finding similarities between data according to the characteristics found in the data and grouping similar data objects into clusters
- Unsupervised learning: NO predefined classes (i.e., learning) by observations vs. learning by examples: supervised)
- Typical applications

1

3

5

- As a stand-alone tool to get insight into data distribution
- As a preprocessing step for other algorithms

Applications Biology: taxonomy of living things: kingdom, phylum, class, order,

Clustering for Data Understanding and

- family, genus and species (界.门.纲.目.科.属.种)
- Information retrieval: documer Land use: Identification of area observation database
- Marketing: Help marketers dis bases, and then use programs
- City-planning: Identify type, value, and geog
- Earth-quake studies: clustered along cont
- Climate: understandi and ocean movemen
- Economic Science: r

Clustering as a Preprocessing Tool (Utility)

- Summarization:
 - Preprocessing for regression, PCA (why?), classification, and association analysis
- Compression:
 - Image processing: vector quantization
- Finding K-nearest Neighbors
 - Localizing search to one or a small number of clusters
- Outlier detection
 - Outliers are often viewed as those "far away" from any cluster

Quality: What Is Good Clustering?

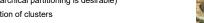
- A good clustering method will produce high quality clusters
 - high intra-class similarity: cohesive within clusters
 - low inter-class similarity: distinctive between clusters
- The quality of a clustering method depends on
 - the similarity measure used by the method
- its implementation, and
- Its ability to discover some or all of the hidden patterns

Measure the Quality of Clustering

- Dissimilarity/Similarity metric
 - Similarity is expressed in terms of a distance function, typically metric: d(i, j)
 - The definitions of distance functions are usually rather different for interval-scaled, boolean, categorical, ordinal ratio, and vector variables
 - Weights should be associated with based on applications and data s
- Quality of clustering:
 - There is usually a separate "qual measures the "goodness" of a clu
 - It is hard to define "similar enough on your enough
 - The answer is typically highly subjective

Considerations for Cluster Analysis

- - Single level vs. hierarchical partitioning (often, multi-level hierarchical partitioning is desirable)



- Exclusive (e.g., one customer belongs to only one region) vs. non exclusive (e.g., one document may belong to more than one class) a.k.a. **multi-label** clustering (e.g., *community detection*)
- - Distance-based (e.g., Euclidian, road network, vector) vs. connectivity-based (e.g., density or contiguity)
- Clustering space

8

Full space (often when low dimensional) vs. subspaces (often in high-dimensional clustering)

7

JOHN B. GOODENOUGH

Requirements and Challenges

- - Clustering all the data instead of only on some samples
- Ability to deal with different types of attributes
 - Numerical, binary, categorical, ordinal, linked, and mixture of
- Constraint-based clustering
 - User may give inputs on constraints
 - Use domain knowledge to determine input parameter
- Interpretability and usability
- Others

9

- Discovery of clusters with arbitrary shapes (e.g., Swiss swirl)
- Ability to deal with noisy data
- Incremental clustering and insensitivity to input order
- High dimensionality



Major Clustering Approaches (I)

- Partitioning approach:
 - Construct various partitions and then evaluate them by some criterion, e.g., minimizing the sum of square errors
 - Typical methods: k-means, k-medoids (PAM), CLARA
- Hierarchical approach:
 - Create a hierarchical decomposition of the set of data (or objects) using some criterion
 - Typical methods: Diana, Agnes, BIRCH, CAMELEON
- Density-based approach:
 - Based on connectivity and density functions
 - Typical methods: DBSACN, OPTICS, DenClue, GraphCut
- Grid-based approach:
 - Based on a multiple-level granularity structure
 - Typical methods: STING, WaveCluster, CLIQUE

10

Major Clustering Approaches (II)

- Model-based:
 - A model is hypothesized for each of the clusters and tries to find the best fit of that model to each other
 - Typical methods: EM, SOM, COBWEB
- Frequent pattern-based:
- Based on the analysis of frequent patterns
- Typical methods: p-Cluster
- User-guided or constraint-based:
 - Clustering by considering user-specified or application-specific constraints
 - Typical methods: COD (obstacles), constrained clustering
- Link-based clustering:
- Objects are often linked together in various ways
- Massive links can be used to cluster objects: SimRank, LinkClus

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Partitioning Algorithms: Basic Concept

Partitioning method: Partitioning a database **D** of **n** objects into a set of ${\it k}$ clusters, such that the sum of squared distances is minimized (where c, is the centroid or medoid of cluster C,)

$$E = \sum_{i=1}^{k} \sum_{p \in C_i} (p - c_i)^2$$

- Given k, find a partition of k clusters that optimizes the chosen partitioning criterion
 - Global optimal: exhaustively enumerate all partitions
 - Heuristic methods: k-means and k-medoids algorithms
 - k-means (MacQueen'67, Lloyd'57/'82): Each cluster is represented
 - k-medoids or PAM (Partition Around Medoids) (Kaufman & Rousseeuw'87): Each cluster is represented by one of the objects in the cluster

The K-Means Clustering Method

- Given k, the k-means algorithm is implemented in four
 - Partition objects into k nonempty subsets
 - Compute seed points as the centroids of the clusters of the current partitioning (the centroid is the center, i.e., mean point, of the cluster)
 - Assign each object to the cluster with the nearest seed point
 - Go back to Step 2, stop when the assignment does not change

Comments on the K-Means Method

Comparing: PAM: O(k(n-k)²), CLARA: O(ks² + k(n-k))

Comment: Often terminates at a local optimum. (initialization can matter!)

Applicable only to objects in a continuous n-dimensional space

Need to specify k, the number of clusters, in advance (there

Not suitable to discover clusters with non-convex shapes

In comparison, k-medoids can be applied to a wider range,

ways to automatically determine the best k (see Hastie et al., 200

Using the k-modes method for categorical data

iterations. Normally, k, t << n.

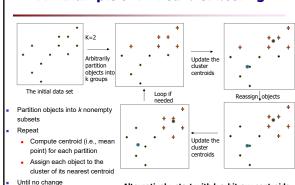
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Strength: Efficient: O(tkn), where n is # objects, k is # clusters, and t is

13

14

An Example of K-Means Clustering



Alternatively, start with k arbitrary centroids

16

Variations of the *K-Means* Method

- Most of the variants of the k-means which differ in
 - Selection of the initial k means
 - Dissimilarity calculations

15

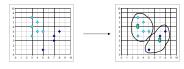
- Strategies to calculate cluster means
- Handling categorical data: k-modes
 - Replacing means of clusters with <u>modes (most popular value)</u>
 - Using new dissimilarity measures to deal with categorical objects
 - Using a <u>frequency</u>-based method to update modes of clusters
 - A mixture of categorical and numerical data: k-prototype method

What Is the Problem of the K-Means Method?

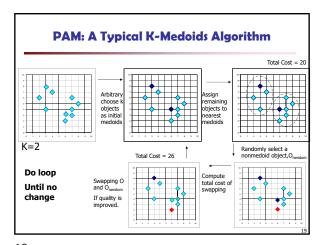
The k-means algorithm is sensitive to outliers!

Sensitive to noisy data and outliers

- Since an object with an extremely large value may substantially distort the distribution of the data (thus the mean calculation)
- K-Medoids: Instead of taking the mean value of the object in a cluster as a reference point, medoids can be used, which is the most centrally located object in a cluster



17 18



The K-Medoid Clustering Method

- K-Medoids Clustering: Find representative objects (medoids) in clusters
 - PAM (Partitioning Around Medoids, Kaufmann & Rousseeuw 1987)
 - Starts from an initial set of medoids and iteratively replaces one
 of the medoids by one of the non-medoids if it improves the total
 distance of the resulting clustering
 - PAM works effectively for small data sets, but <u>does not scale</u>
 well for large data sets (due to the computational complexity)
- Efficiency improvement on PAM
 - CLARA (Kaufmann & Rousseeuw, 1990): PAM on samples
 - CLARANS (Ng & Han, 1994): Randomized re-sampling

19

20



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Cluster Analysis: Basic Concepts

Partitioning Methods

Hierarchical Methods 🦊

Density-Based Methods

Grid-Based Methods

Evaluation of Clustering

Summary

HW1-3 Graded before midterm

21

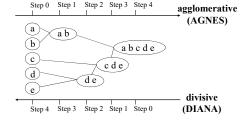
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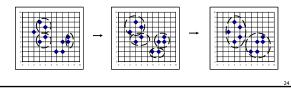
Hierarchical Clustering

 Use distance matrix as clustering criteria. This method does not require the number of clusters k as an input, but needs a termination condition



AGNES (Agglomerative Nesting)

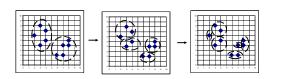
- Introduced in Kaufmann and Rousseeuw (1990)
- Implemented in statistical packages, e.g., Splus
- Use the single-link method and the dissimilarity matrix
- Merge nodes that have the least dissimilarity
- Go on in a non-descending fashion
- Eventually all nodes belong to the same cluster



Decompose data objects into several levels of nested partitioning (tree of clusters), called a dendrogram A clustering of the data objects is obtained by cutting the dendrogram at the desired level, then each connected component forms a cluster

DIANA (Divisive Analysis)

- Introduced in Kaufmann and Rousseeuw (1990)
- Implemented in statistical analysis packages, e.g., Splus
- Inverse order of AGNES
- Eventually each node forms a cluster on its own



26

Distance between Clusters

25

27

29





- Single link: smallest distance between an element in one cluster and an element in the other, i.e., dist(K_i, K_j) = min(t_{ip}, t_{iq})
- $\label{eq:complete link: largest distance between an element in one cluster and an element in the other, i.e., \ dist(K_i, K_j) = max(t_{ip}, t_{jq})$
- Average: avg distance between an element in one cluster and an element in the other, i.e., $dist(K_i, K_j) = avg(t_{ip}, t_{jq})$
- Centroid: distance between the centroids of two clusters, i.e., dist(K_i, K_j) = dist(C_i, C_j)
- $\begin{tabular}{ll} \hline & Medold: distance between the medolds of two clusters, i.e., & dist(K_i, K_j) = dist(M_i, M_j) \\ \hline \end{tabular}$
 - Medoid: a chosen, <u>centrally located</u> object in the cluster

Centroid, Radius and Diameter of a Cluster (for numerical data sets)

Centroid: the "middle" of a cluster

$$C_m = \frac{\sum_{i=1}^{N} (t_{ip})}{N}$$

- Radius: square root of average distance from any point of the cluster to its **centroid** $R_m = \sqrt{\frac{\sum_{i=1}^N (t_{ip} c_m)^2}{\sum_{i=1}^N (t_{ip} c_m)^2}}$
- Diameter: square root of average mean squared distance between all pairs of points in the cluster

$$D_{m} = \sqrt{\frac{\sum_{i=1}^{N} \sum_{i=1}^{N} (t_{ip} - t_{iq})^{2}}{N(N-1)}}$$

28

30

Extensions to Hierarchical Clustering

- Major weakness of agglomerative clustering methods
 - Can never undo what was done previously
 - <u>Do not scale</u> well: time complexity of at least O(n²),
 where n is the number of total objects
- Integration of hierarchical & distance-based clustering
 - BIRCH (1996): uses CF-tree and incrementally adjusts the quality of sub-clusters
 - CHAMELEON (1999): hierarchical clustering using dynamic modeling

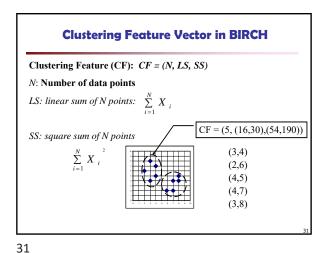
BIRCH (Balanced Iterative Reducing and Clustering Using Hierarchies)

Zhang, Ramakrishnan & Livny, SIGMOD'96



- Incrementally construct a CF (Clustering Feature) tree, a hierarchical data structure for multiphase clustering
 - Phase 1: scan DB to build an initial in-memory CF tree (a multi-level compression of the data that tries to preserve the inherent clustering structure of the data)
 - Phase 2: use an arbitrary clustering algorithm to cluster the leaf nodes of the CF-tree
- Scales linearly: finds a good clustering with a single scan and improves the quality with a few additional scans
- Weakness: handles only numeric data, and <u>sensitive to the order</u> of the data record

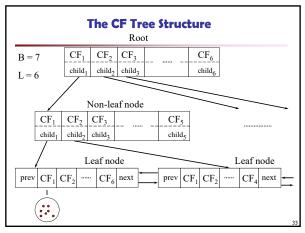
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CF-Tree in BIRCH

- Clustering feature:
 - Summary of the statistics for a given subcluster: the 0-th, 1st, and 2nd moments of the subcluster from the statistical point
 - Registers crucial measurements for computing cluster and utilizes storage efficiently
- A CF tree is a height-balanced tree that stores the clustering features for a hierarchical clustering
 - A nonleaf node in a tree has descendants or "children"
 - The nonleaf nodes store **sums** of the CFs of their children
- A CF tree has two parameters
 - Branching factor: max # of children
 - Threshold: max diameter of sub-clusters stored at the leaf

32



The Birch Algorithm

Cluster Diameter

$$\sqrt{\frac{1}{n(n-1)}\sum \left(x_i-x_j\right)^2}$$

- For each point in the input
 - Find closest leaf entry
 - Add point to leaf entry and update CF
 - If entry diameter > max_diameter, then split leaf, and possibly parents
- Algorithm is O(n)
- Concerns

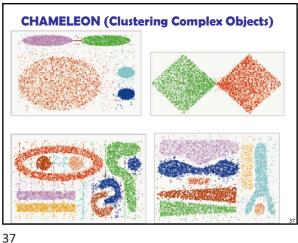
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- Sensitive to insertion order of data points
- Since we fix the size of leaf nodes, so clusters may not be so natural
- Clusters tend to be spherical given the radius and diameter

33

- **CHAMELEON: Hierarchical Clustering Using Dynamic Modeling (1999)**
- CHAMELEON: G. Karypis, E. H. Han, and V. Kumar, 1999
- Measures the similarity based on a dynamic model
 - Two clusters are merged only if the interconnectivity and closeness (proximity) between two clusters are high relative to the internal interconnectivity of the clusters and closeness of items within the clusters
- Graph-based, and a two-phase algorithm
 - 1. Use a graph-partitioning algorithm: cluster objects into a large number of relatively small sub-clusters
 - 2. Use an agglomerative hierarchical clustering algorithm: find the genuine clusters by repeatedly combining these sub-clusters

Overall Framework of CHAMELEON Sparse Graph K-NN Graph P and q are <u>connected</u> if q is among the top k closest neighbors of p Relative interconnectivity connectivity of c₁ and c₂ over internal connectivity Relative closeness: closeness of c₁ and c₂ ov internal closeness



Generative Model

• Given a set of 1-D points $X = \{x_1, ..., x_n\}$ for clustering analysis & assuming they are generated by a Gaussian distribution:

 $\mathcal{N}(\mu, \sigma^2) = \cdot$

• The probability that a point $x_i \in X$ is generated by the

 $P(x_i|\mu,\sigma^2) = \frac{1}{\sqrt{2\pi\sigma^2}}e^{-\frac{1}{2\sigma^2}}$

• The likelihood that X is generated by the model:

 $L(N(\mu, \sigma^2) : X) = P(X|\mu, \sigma^2) = \prod_{i=1}^{n} \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{(x_i - \mu)^2}{2\sigma^2}}$

The task of learning the generative model: find the parameters μ and σ^2 such that the maximum likelihood

 $\mathcal{N}(\mu_0, \sigma_0^2) = \arg\max\{L(\mathcal{N}(\mu, \sigma^2) : X)\}$

39

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Probabilistic Hierarchical Clustering

- Algorithmic hierarchical clustering
 - Nontrivial to choose a good distance measure
 - Hard to handle missing attribute values
 - Optimization goal not clear: heuristic, local search
- Probabilistic hierarchical clustering
- Use probabilistic models to measure distances between clusters
- Generative model: Regard the set of data objects to be clustered as a sample of the underlying data generation mechanism to be analyzed
- Easy to understand, same efficiency as algorithmic agglomerative clustering method, can handle partially observed data
- In practice, assume the generative models adopt common distributions functions, e.g., Gaussian distribution or Bernoulli distribution, governed by parameters

38

A Probabilistic Hierarchical Clustering Algorithm

For a set of objects partitioned into m clusters C_1, \ldots, C_m the quality can be measured by, $Q(\{C_1,\dots,C_m\}) = \prod_{i=1}^m P(C_i)$

where P() is the maximum likelihood

- $\begin{array}{ll} \textbf{Distance} \ \ \text{between clusters} \ \ C_1 \ \text{and} \ \ C_2 : \ \ dist(C_i,C_j) = -\log \frac{P(C_1 \cup C_2)}{P(C_1)P(C_2)} \\ \textbf{Algorithm:} \ \ \text{Progressively merge points and clusters} \end{array}$
- Input: $D = \{o_1, ..., o_n\}$: a data set containing n objects Output: A hierarchy of clusters Method

Create a cluster for each object $C_i = \{o_i\}, 1 \le i \le n$; For i = 1 to n {

Find pair of clusters C_i and C_i such that $C_i, C_j = \operatorname{argmax}_{i \neq j} \{ \log (P(C_i \cup C_j)/(P(C_i)P(C_j)) \};$

If log $(P(C_i \cup C_i)/(P(C_i)P(C_i)) > 0$ then merge C_i and C_i

40

Density-Based Clustering Methods

- Clustering based on density (local cluster criterion), such as *density-connected* points
- Major features:
 - Discover clusters of arbitrary shape
 - Handle noise
 - One scan
 - Need density parameters as termination condition
- Several interesting studies:
 - DBSCAN: Ester, et al. (KDD'96)
 - OPTICS: Ankerst, et al (SIGMOD'99).
 - <u>DENCLUE</u>: Hinneburg & D. Keim (KDD'98)
 - <u>CLIQUE</u>: Agrawal, et al. (SIGMOD'98) (more grid-based)

Density-Based Clustering: Basic Concepts

- Two parameters:
 - **Eps**: Maximum radius of the neighbourhood
 - MinPts: Minimum number of points in an Epsneighbourhood of that point
- $N_{Eps}(p)$: {q belongs to D | dist(p,q) \leq Eps}
- Directly density-reachable: A point p is directly density-reachable from a point q w.r.t. Eps, MinPts if
 - p belongs to N_{Eps}(q)
 - core point condition:

 $|N_{Eps}(q)| \ge MinPts$



MinPts = 5

Eps = 1 cm

Density-Reachable and Density-Connected

- Density-reachable:
 - A point p is density-reachable from a point q w.r.t. Eps, MinPts if there is a chain of points $p_1, ..., p_n, p_1 = q$, $p_n = p$ such that $\underline{p_{i+1}}$ is directly density-reachable from $\underline{p_i}$



- Density-connected
 - A point p is density-connected to a point q w.r.t. Eps, MinPts if there is a point o such that both, p and q are density-reachable from o w.r.t. Eps and MinPts

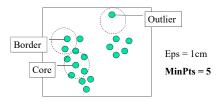


43

44

DBSCAN: Density-Based Spatial Clustering of Applications with Noise

- Relies on a density-based notion of cluster: A cluster is defined as a maximal set of density-connected points
- Discovers clusters of arbitrary shape in spatial databases with noise



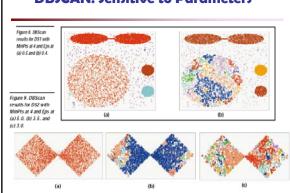
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DBSCAN: The Algorithm

- Arbitrary select a point p
- Retrieve all points (directly*) density-reachable from p w.r.t. Eps and MinPts
- If p is a core point, a cluster is formed
- If p is a border point*, (or) no points are densityreachable from p and DBSCAN visits the next point of the database
- Continue the process until all of the points have been processed

46

DBSCAN: Sensitive to Parameters



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 OPTICS: Ordering Points To Identify the Clustering Structure

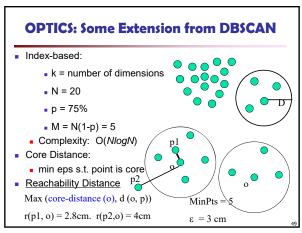
Ankerst, Breunig, Kriegel, and Sander (SIGMOD'99)

OPTICS: A Cluster-Ordering Method (1999)

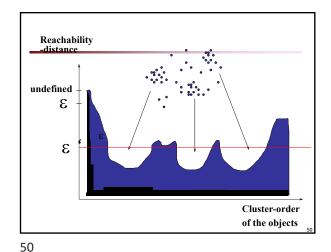
- Produces a special order of the database wrt its density-based clustering structure
- This cluster-ordering contains info equiv. to the densitybased clusterings corresponding to a broad range of parameter settings (s.t. insensitive to parameters)
- Good for both automatic and interactive cluster analysis, including finding intrinsic clustering structure
- Can be represented graphically or using visualization techniques

48

47 48



49



Density-Based Clustering: OPTICS & Its Applications

DENCLUE: Using Statistical Density Functions

DENsity-based CLUstEring by Hinneburg & Keim (KDD'98)
Using statistical density functions: $f_{Gaussian}(x,y) = e^{\frac{d(x,y)^2}{2\sigma^2}} \qquad f_{Gaussian}^D(x,x_i) = \sum_{i=1}^N e^{-\frac{d(x,x_i)^2}{2\sigma^2}}$ Major features

Solid mathematical foundation
Good for data sets with large amounts of noise
Allows a compact mathematical description of arbitrarily shaped clusters in high-dimensional data sets (i.e., mixture of Gaussian)
Significantly faster than existing algorithm (e.g., DBSCAN)
But needs a large number of parameters

52

51

Denclue: Technical Essence

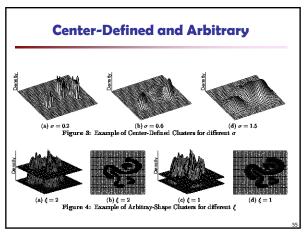
- Uses grid cells but only keeps information about grid cells that do actually contain data points and manages these cells in a tree-based access structure
- Influence function: describes the impact of a data point within its neighborhood
- Overall density of the data space can be calculated as the sum of the influence function of all data points
- Clusters can be determined mathematically by identifying density attractors
- <u>Density attractors</u> are local maximal of the overall density function
- Center defined clusters: assign to each density attractor the points density attracted to it
- <u>Arbitrary shaped cluster</u>: merge density attractors that are connected through paths of high density (> threshold)

Density Attractor

(a) Data Set
(c) Gaussian

Data Space

53 54



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55 56

Grid-Based Clustering Method

- Using <u>multi-resolution grid data structure</u>
- Several interesting methods

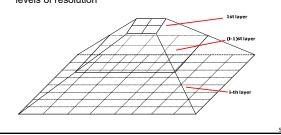
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59

- STING (a STatistical Information Grid approach) by Wang, Yang and Muntz (1997)
- WaveCluster by Sheikholeslami, Chatterjee, and Zhang (VLDB'98)
 - A multi-resolution clustering approach using wavelet method
- CLIQUE: Agrawal, et al. (SIGMOD'98)
 - Both grid-based and subspace clustering

STING: A Statistical Information Grid Approach

- Wang, Yang and Muntz (VLDB'97)
- The spatial area is divided into rectangular cells
- There are several <u>levels</u> of cells corresponding to different levels of resolution



58

60

The STING Clustering Method

- Each cell at a high level is partitioned into a number of smaller cells in the next lower level
- Statistical info of each cell is calculated and stored beforehand and is used to answer queries
- Parameters of higher level cells can be easily calculated from parameters of lower level cell
 - count, mean, s, min, max
 - type of distribution—normal, uniform, etc.
- Use a top-down approach to answer spatial data queries
- Start from a pre-selected layer—typically with a small number of cells
- For each cell in the current level compute the confidence interval

STING Algorithm and Its Analysis

- Remove the irrelevant cells from further consideration
- When finish examining the current layer, proceed to the next lower level
- Repeat this process until the bottom layer is reached
- Advantages:
 - Query-independent, easy to parallelize, incremental update
 - $\,\blacksquare\,\, {\it O(K)},$ where ${\it K}$ is the number of grid cells at the lowest level
- Disadvantages:
 - All the <u>cluster boundaries</u> are either horizontal or vertical, and no diagonal boundary is detected

CLIQUE (Clustering In QUEst)

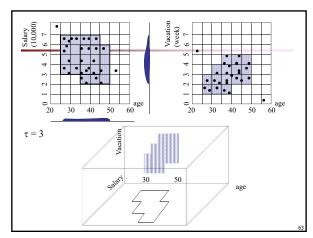
- Agrawal, Gehrke, Gunopulos, Raghavan (SIGMOD'98)
- Automatically identifying <u>subspaces</u> of a high dimensional data space that allow better clustering than original space
- CLIQUE can be considered as <u>both</u> density-based and grid-based
 - It partitions each dimension into the same number of equal length into rel.
 - It partitions an m-dimensional data space into non-overlapping rectangular units
 - A unit is dense if the fraction of total data points contained in the unit exceeds the input model parameter
 - A cluster is a maximal set of connected dense units within a subspace

CLIQUE: The Major Steps

- Partition the data space and find the number of points that lie inside each cell of the partition.
- Identify the subspaces that contain clusters using the Apriori principle
- Identify clusters
 - Determine dense units in all subspaces of interests
 - Determine connected dense units in all subspaces of interests
- Generate minimal description for the clusters
 - Determine maximal regions that cover a cluster of connected dense units for each cluster
 - Determination of minimal cover for each cluster

61

62



Strength and Weakness of CLIQUE

Strength

- automatically finds subspaces of the highest dimensionality such that high density clusters exist in those subspaces
- insensitive to the order of records in input and does not presume some canonical data distribution
- scales linearly with the size of input and has good scalability as the number of dimensions in the data increases
- Weaknes
 - The accuracy of the clustering result may be degraded at the expense of simplicity of the method

63

64

Chapter 10. Cluster Analysis: Basic Concepts and Methods

- Cluster Analysis: Basic Concepts
- Partitioning Methods
- Hierarchical Methods
- Density-Based Methods
- Grid-Based Methods
- Evaluation of Clustering
- Summary

Assessing Clustering Tendency

- Assess if non-random structure exists in the data by measuring the probability that the data is generated by a *uniform* data distribution
- Test spatial randomness by statistic test: Hopkins Static
- Given a dataset D regarded as a sample of a random variable o, determine how far away o is from being uniformly distributed in the data space
- Sample n points, p_1, \ldots, p_n uniformly from D. For each p_i , find its nearest neighbor in D: $x_i = \min\{dist\ (p_i, v)\}$ where v in D
- Sample n points, q₁, ..., q_n uniformly from D. For each q_n find its nearest neighbor in D − {q_n}: y_i = min{dist (q_n v)} where v in D and v ≠ q_i
- v ≠ q_i Calculate the Hopkins Statistic: $H = \frac{\sum_{i=1}^n y_i}{\sum_{i=1}^n x_i + \sum_{i=1}^n y_i}$
- If D is uniformly distributed, $\sum x_i$ and $\sum y_i$ will be close to each other and H is close to 0.5. If D is highly skewed, H is close to 0

6

65

Determine the Number of Clusters

- Empirical method
 - # of clusters ≈√n/2 for a dataset of n points
- Elbow method
 - Use the turning point in the curve of sum of within cluster variance w.r.t the # of clusters
- Cross validation method
 - Divide a given data set into m parts
 - Use m 1 parts to obtain a clustering model
 - Use the remaining part to test the quality of the clustering
 - E.g., For each point in the test set, find the closest centroid, and use the sum of squared distance between all points in the test set and the closest centroids to measure how well the model fits the test set
 - For any k > 0, repeat it m times, compare the overall quality measure w.r.t. different k's, and find # of clusters that fits the data the best

Measuring Clustering Quality

- Two methods: extrinsic vs. intrinsic
- Extrinsic: supervised, i.e., the ground truth is available
 - Compare a clustering against the ground truth using certain clustering quality measure
 - Ex. BCubed precision and recall metrics
- Intrinsic: unsupervised, i.e., the ground truth is unavailable
 - Evaluate the goodness of a clustering by considering how well the clusters are separated, and how compact the clusters are
 - Ex. Silhouette coefficient, BIC (Bayesian Information Criterion), AIC (Akaike information criterion)

7

67

Measuring Clustering Quality: Extrinsic Methods

- Clustering quality measure: $Q(C, C_g)$, for a clustering C given the ground truth $C_{g\cdot}$
- Q is good if it satisfies the following 4 essential criteria
 - Cluster homogeneity: the purer, the better
 - Cluster completeness: should assign objects belong to the same category in the ground truth to the same cluster.
 - Rag bag: putting a heterogeneous object into a pure cluster should be penalized more than putting it into a rag bag (i.e., "miscellaneous" or "other" category)
 - Small cluster preservation: splitting a small category into pieces is more harmful than splitting a large category into pieces

68

70

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69

Summary

- Cluster analysis groups objects based on their similarity and has wide applications
- Measure of similarity can be computed for various types of data
- Clustering algorithms can be categorized into partitioning methods, hierarchical methods, density-based methods, grid-based methods, and model-based methods
- K-means and K-medoids algorithms are popular partitioning-based clustering algorithms
- Birch and Chameleon are interesting hierarchical clustering algorithms, and there are also probabilistic hierarchical clustering algorithms
- DBSCAN, OPTICS, and DENCLU are interesting density-based algorithms
- STING and CLIQUE are grid-based methods, where CLIQUE is also a subspace clustering algorithm
- Quality of clustering results can be evaluated in various ways

Homework Assignment #5

Textbook

- **1**0.2, 10.4, 10.6
- More after the midterm review
- May include using some toolboxes

71

72

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74

75

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